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<!--StartFragment-->RESULT 13
US-09-006-841-4
; Sequence 4, Application US/09006841
; Patent No. 6468798
; GENERAL INFORMATION:
;   APPLICANT: Robert J. Debs
;   TITLE OF INVENTION: EXPRESSION OF CLONED
;   TITLE OF INVENTION: GENES IN THE
;   TITLE OF INVENTION: LUNG BY AEROSOL - AND
;   TITLE OF INVENTION: LIPOSOME-BASED
;   TITLE OF INVENTION: DELIVERY
;   NUMBER OF SEQUENCES: 4
;   CORRESPONDENCE ADDRESS:
;     ADDRESSEE: Fish & Richardson
;     STREET: 2200 Sand Hill Road,
;     STREET: Suite 100
;     CITY: Menlo Park
;     STATE: California
;     COUNTRY: U.S.A.
;     ZIP: 94025
;   COMPUTER READABLE FORM:
;     MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
;     COMPUTER: IBM PS/2 Model 50Z or 55SX
;     OPERATING SYSTEM: MS-DOS (Version 5.0)
;     SOFTWARE: WordPerfect (Version 5.1)
;   CURRENT APPLICATION DATA:
;     APPLICATION NUMBER: US/09/006,841
;     FILING DATE:
;     CLASSIFICATION:
;   PRIOR APPLICATION DATA:
;     APPLICATION NUMBER: US/07/972,135
;     FILING DATE: No. 6468798ember 5, 1992
;     APPLICATION NUMBER: 07/809,291
;     FILING DATE: December 17, 1991
;   ATTORNEY/AGENT INFORMATION:
;     NAME: Barbara Rae-Venter
;     REGISTRATION NUMBER: 32,750
;     REFERENCE/DOCKET NUMBER: 05935/007US1
;   TELECOMMUNICATION INFORMATION:
;     TELEPHONE: (415) 854-5277
;     TELEFAX: (415) 854-0875
;     TELEX: 200154
;   INFORMATION FOR SEQ ID NO: 4:
;     SEQUENCE CHARACTERISTICS:
;       LENGTH: 930
;       TYPE: nucleic acid
;       STRANDEDNESS: single
;       TOPOLOGY: linear
US-09-006-841-4

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Query Match          100.0%; Score 366; DB 3; Length 930;
Best Local Similarity 100.0%; Pred. No. 1.4e-116;
Matches 366; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Db      156 TAGTTATTAATAGTAATCAATTACGGGGTCATTAGTTCATAGCCCATATATGGAGTTC CG 215

Qy      61 CGTTACATAACTTACGGTAAATGGCCCGCTGGCTGACCGCCCAACGACCCCCGCCCAT 120
        |||
Db      216 CGTTACATAACTTACGGTAAATGGCCCGCTGGCTGACCGCCCAACGACCCCCGCCCAT 275

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Qy      181  ATGGGTGGAGTATTTACGGTAAACTGCCCACTTGGCAGTACATCAAGTGTATCATATGCC 240
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      336  ATGGGTGGAGTATTTACGGTAAACTGCCCACTTGGCAGTACATCAAGTGTATCATATGCC 395

Qy      241  AAGTACGCCCCCTATTGACGTCAATGACGGTAAATGGCCCGCCTGGCATTATGCCCAGTA 300
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Qy      301  CATGACCTTATGGGACTTTCCTACTTGGCAGTACATCTACGTATTAGTCATCGCTATTAC 360
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Qy      361  CATGGT 366
        |||||
Db      516  CATGGT 521
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